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Sequence Listing was accepted.
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Reviewer: markspencer
Timestamp: [year=2008; month=11; day=25; hr=8; min=29; sec=34; ms=74;]
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Validated By CRFValidator v 1.0.3

Application No: 10511384 Version No: 4.0

Input Set:

Output Set:

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Finished: 2008-10-31 15:17:08.975
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 195 ms
Total Warnings: 218
Total Errors: 0
No. of SeqIDs Defined: 226
Actual SeqID Count: 226

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Input Set:

Output Set:

Started: 2008-10-31 15:17:06.780
Finished: 2008-10-31 15:17:08.975
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 195 ms
Total Warnings: 218
Total Errors: 0
No. of SeqIDs Defined: 226
Actual SeqID Count: 226

Error code	Error Description

This error has occured more than 20 times, will not be displayed

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<120> Antiangiogenic active immunotherapies

<130> 976-19 PCT/US/RCE

<140> 10511384
<141> 2008-10-31

<150> CU 2002/0076
<151> 2002-04-15

<160> 226

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<212> PRT
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<400> 147

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Tyr Leu His His Ala Lys Trp Ser Gln Ala Ala Pro Met Ala Glu Gly
20 25 30

Gly Gly Gln Asn His His Glu Val Val Lys Phe Met Asp Val Tyr Gln
35 40 45

Arg Ser Tyr Cys His Pro Ile Glu Thr Leu Val Asp Ile Phe Gln Glu
50 55 60

Tyr Pro Asp Glu Ile Glu Tyr Ile Phe Lys Pro Ser Cys Val Pro Leu
65 70 75 80

Met Arg Cys Gly Gly Cys Cys Asn Asp Glu Gly Leu Glu Cys Val Pro
85 90 95

Thr Glu Glu Ser Asn Ile Thr Met Gln Ile Met Glu Ile Glu Pro Glu
100 105 110

Gln Gly Gln His Ile Gly Glu Met Ser Phe Leu Gln His Asn Lys Cys
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Pro Arg Arg
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<211> 444
<212> DNA
<213> Homo Sapiens

<400> 20

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gtgaagttca tggatgtcta tcagcgcagc tactgccatc caatcgagac cctggtggac 180

atcttccagg agtaccctga tgagatcgag tacatcttca agccatcctg tgtgcccctg 240

atgcgatgcg ggggctgctg caatgacgag ggcctggagt gtgtgcccac tgaggagtcc 300

aacatcacca tgcagattat gcggatcaaa cctcaccaag gccagcacat aggagagatg 360

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<211> 147
<212> PRT
<213> Homo Sapiens

<400> 147

Met Asn Phe Leu Leu Ser Trp Val His Trp Ser Leu Ala Leu Leu Leu
1 5 10 15

Tyr Leu His His Ala Lys Trp Ser Gln Ala Ala Pro Met Ala Glu Gly
20 25 30

Gly Gly Gln Asn His His Glu Val Val Lys Phe Met Asp Val Tyr Gln
35 40 45

Arg Ser Tyr Cys His Pro Ile Glu Thr Leu Val Asp Ile Phe Gln Glu
50 55 60

Tyr Pro Asp Glu Ile Glu Tyr Ile Phe Lys Pro Ser Cys Val Pro Leu
65 70 75 80

Met Arg Cys Gly Gly Cys Cys Asn Asp Glu Gly Leu Glu Cys Val Pro
85 90 95

Thr Glu Glu Ser Asn Ile Thr Met Gln Ile Met Ala Ile Ala Pro Ala
100 105 110

Gln Gly Gln His Ile Gly Glu Met Ser Phe Leu Gln His Asn Lys Cys
115 120 125

Glu Cys Arg Pro Lys Lys Asp Arg Ala Arg Gln Glu Lys Cys Asp Lys
130 135 140

Pro Arg Arg
145

<210> 22
<211> 444
<212> DNA
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atcttccagg agtaccctga tgagatcgag tacatcttca agccatcctg tgtgcccctg    240

atgcgatgcg ggggctgctg caatgacgag ggcctggagt gtgtgcccac tgaggagtcc    300

aacatcacca tgcagattat ggcaatcgca cctgcacaag gccagcacat aggagagatg    360

agcttcctac agcacaacaa atgtgaatgc agaccaaaga aagatagagc aagacaagaa    420

aaatgtgaca agccgaggcg gtaa                                           444
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<210> 23
<211> 314
<212> PRT
<213> Homo Sapiens

<400> 314

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Met Glu Ser Lys Val Leu Leu Ala Val Ala Leu Trp Leu Cys Val Glu
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Thr Arg Ala Ala Ser Val Gly Leu Pro Ser Val Ser Leu Asp Leu Pro
             20                  25                  30

Arg Leu Ser Ile Gln Lys Asp Ile Leu Thr Ile Lys Ala Asn Thr Thr
         35                  40                  45

Leu Gln Ile Thr Cys Arg Gly Gln Arg Asp Leu Asp Trp Leu Trp Pro
     50                  55                  60

Asn Asn Gln Ser Gly Ser Glu Gln Arg Val Glu Val Thr Glu Cys Ser
 65                  70                  75                  80

Asp Gly Leu Phe Cys Lys Thr Leu Thr Ile Pro Lys Val Ile Gly Asn
                 85                  90                  95

Asp Thr Gly Ala Tyr Lys Cys Phe Tyr Arg Glu Thr Asp Leu Ala Ser
            100                 105                 110

Val Ile Tyr Val Tyr Val Gln Asp Tyr Arg Ser Pro Phe Ile Ala Ser
        115                 120                 125

Val Ser Asp Gln His Gly Val Val Tyr Ile Thr Glu Asn Lys Asn Lys
    130                 135                 140

Thr Val Val Ile Pro Cys Leu Gly Ser Ile Ser Asn Leu Asn Val Ser
145                 150                 155                 160

Leu Cys Ala Arg Tyr Pro Glu Lys Arg Phe Val Pro Asp Gly Asn Arg
                165                 170                 175
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Ile Ser Trp Asp Ser Lys Lys Gly Phe Thr Ile Pro Ser Tyr Met Ile
            180                 185                 190

Ser Tyr Ala Gly Met Val Phe Cys Glu Ala Lys Ile Asn Asp Glu Ser
        195                 200                 205

Tyr Gln Ser Ile Met Tyr Ile Val Val Val Val Gly Tyr Arg Ile Tyr
    210                 215                 220

Asp Val Val Leu Ser Pro Ser His Gly Ile Glu Leu Ser Val Gly Glu
225                 230                 235                 240

Lys Leu Val Leu Asn Cys Thr Ala Arg Thr Glu Leu Asn Val Gly Ile
                245                 250                 255

Asp Phe Asn Trp Glu Tyr Pro Ser Ser Lys His Gln His Lys Lys Leu
            260                 265                 270

Val Asn Arg Asp Leu Lys Thr Gln Ser Gly Ser Glu Met Lys Lys Phe
        275                 280                 285

Leu Ser Thr Leu Thr Ile Asp Gly Val Thr Arg Ser Asp Gln Gly Leu
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Tyr Thr Cys Ala Ala Ser Ser Gly Leu Met
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<210> 24
<211> 943
<212> DNA
<213> Homo Sapiens

<400> 24

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cttacaatta aggctaatac aactcttcaa attacttgca ggggacagag ggacttggac    180

tggctttggc ccaataatca gagtggcagt gagcaaaggg tggaggtgac tgagtgcagc    240

gatggcctct tctgtaagac actcacaatt ccaaaagtga tcggaaatga cactggagcc    300

tacaagtgct tctaccggga aactgacttg gcctcggtca tttatgtcta tgttcaagat    360

tacagatctc catttattgc ttctgttagt gaccaacatg gagtcgtgta cattactgag    420

aacaaaaaca aaactgtggt gattccatgt ctcgggtcca tttcaaatct caacgtgtca    480

ctttgtgcaa gatacccaga aaagagattt gttcctgatg gtaacagaat ttcctgggac    540

agcaagaagg gctttactat tcccagctac atgatcagct atgctggcat ggtcttctgt    600

gaagcaaaaa ttaatgatga aagttaccag tctattatgt acatagttgt cgttgtaggg    660
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tataggattt atgatgtggt tctgagtccg tctcatggaa ttgaactatc tgttggagaa 720

aagcttgtct taaattgtac agcaagaact gaactaaatg tggggattga cttcaactgg 780

gaataccctt cttcgaagca tcagcataag aaacttgtaa accgagacct aaaaacccag 840

tctgggagtg agatgaagaa atttttgagc accttaacta tagatggtgt aacccggagt 900

gaccaaggat tgtacacctg tgcagcatcc agtgggctga tga 943

<210> 25
<211> 611
<212> PRT
<213> Homo sapiens

<400> 611

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Asn Leu Glu Ile Ile Ile Leu Val Gly Thr Ala Val Ile Ala Met Phe
20 25 30

Phe Trp Leu Leu Leu Val Ile Ile Leu Arg Thr Val Lys Arg Ala Asn
35 40 45

Gly Gly Glu Leu Lys Thr Gly Tyr Leu Ser Ile Val Met Asp Pro Asp
50 55 60

Glu Leu Pro Leu Asp Glu His Cys Glu Arg Leu Pro Tyr Asp Ala Ser
65 70 75 80

Lys Trp Glu Phe Pro Arg Asp Arg Leu Lys Leu Gly Lys Pro Leu Gly
85 90 95

Arg Gly Ala Phe Gly Gln Val Ile Glu Ala Asp Ala Phe Gly Ile Asp
100 105 110

Lys Thr Ala Thr Cys Arg Thr Val Ala Val Lys Met Leu Lys Glu Gly
115 120 125

Ala Thr His Ser Glu His Arg Ala Leu Met Ser Glu Leu Lys Ile Leu
130 135 140

Ile His Ile Gly His His Leu Asn Val Val Asn Leu Leu Gly Ala Cys
145 150 155 160

```
Thr Lys Pro Gly Gly Pro Leu Met Val Ile Val Glu Phe Cys Lys Phe
                165                 170                 175

Gly Asn Leu Ser Thr Tyr Leu Arg Ser Lys Arg Asn Glu Phe Val Pro
            180                 185                 190

Tyr Lys Thr Lys Gly Ala Arg Phe Arg Gln Gly Lys Asp Tyr Val Gly
        195                 200                 205

Ala Ile Pro Val Asp Leu Lys Arg Arg Leu Asp Ser Ile Thr Ser Ser
    210                 215                 220

Gln Ser Ser Ala Ser Ser Gly Phe Val Glu Glu Lys Ser Leu Ser Asp
225                 230                 235                 240

Val Glu Glu Glu Glu Ala Pro Glu Asp Leu Tyr Lys Asp Phe Leu Thr
                245                 250                 255

Leu Glu His Leu Ile Cys Tyr Ser Phe Gln Val Ala Lys Gly Met Glu
            260                 265                 270

Phe Leu Ala Ser Arg Lys Cys Ile His Arg Asp Leu Ala Ala Arg Asn
        275                 280                 285

Ile Leu Leu Ser Glu Lys Asn Val Val Lys Ile Cys Asp Phe Gly Leu
    290                 295                 300

Ala Arg Asp Ile Tyr Lys Asp Pro Asp Tyr Val Arg Lys Gly Asp Ala
305                 310                 315                 320

Arg Leu Pro Leu Lys Trp Met Ala Pro Glu Thr Ile Phe Asp Arg Val
                325                 330                 335

Tyr Thr Ile Gln Ser Asp Val Trp Ser Phe Gly Val Leu Leu Trp Glu
            340                 345                 350

Ile Phe Ser Leu Gly Ala Ser Pro Tyr Pro Gly Val Lys Ile Asp Glu
        355                 360                 365

Glu Phe Cys Arg Arg Leu Lys Glu Gly Thr Arg Met Arg Ala Pro Asp
    370                 375                 380
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Tyr Thr Thr Pro Glu Met Tyr Gln Thr Met Leu Asp Cys Trp His Gly
385                 390                 395                 400

Glu Pro Ser Gln Arg Pro Thr Phe Ser Glu Leu Val Glu His Leu Gly
                405                 410                 415

Asn Leu Leu Gln Ala Asn Ala Gln Gln Asp Gly Lys Asp Tyr Ile Val
            420                 425                 430

Leu Pro Ile Ser Glu Thr Leu Ser Met Glu Glu Asp Ser Gly Leu Ser
        435                 440                 445

Leu Pro Thr Ser Pro Val Ser Cys Met Glu Glu Glu Glu Val Cys Asp
    450                 455                 460

Pro Lys Phe His Tyr Asp Asn Thr Ala Gly Ile Ser Gln Tyr Leu Gln
465                 470                 475                 480

Asn Ser Lys Arg Lys Ser Arg Pro Val Ser Val Lys Thr Phe Glu Asp
                485                 490                 495

Ile Pro Leu Glu Glu Pro Glu Val Lys Val Ile Pro Asp Asp Asn Gln
            500                 505                 510

Thr Asp Ser Gly Met Val Leu Ala Ser Glu Glu Leu Lys Thr Leu Glu
        515                 520                 525

Asp Arg Thr Lys Leu Ser Pro Ser Phe Gly Gly Met Val Pro Ser Lys
    530                 535                 540

Ser Arg Glu Ser Val Ala Ser Glu Gly Ser Asn Gln Thr Ser Gly Tyr
545                 550                 555                 560

Gln Ser Gly Tyr His Ser Asp Asp Thr Asp Thr Thr Val Tyr Ser Ser
                565                 570                 575

Glu Glu Ala Glu Leu Leu Lys Leu Ile Glu Ile Gly Val Gln Thr Gly
            580                 585                 590

Ser Thr Ala Gln Ile Leu Gln Pro Asp Ser Gly Thr Thr Leu Ser Ser
        595                 600                 605

Pro Pro Val
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610

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<212> DNA
<213> Homo Sapiens

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ctacggaccg ttaagcgggc caatggaggg gaactgaaga caggctactt gtccatcgtc 180

atggatccag atgaactccc attggatgaa cattgtgaac gactgcctta tgatgccagc 240

aaatgggaat tccccagaga ccggctgaag ctaggtaagc ctcttggccg tggtgccttt 300

ggccaagtga ttgaagcaga tgcctttgga attgacaaga cagcaacttg caggacagta 360

gcagtcaaaa tgttgaaaga aggagcaaca cacagtgagc atcgagctct catgtctgaa 420

ctcaagatcc tcattcatat tggtcaccat ctcaatgtgg tcaaccttct aggtgcctgt 480

accaagccag gagggccact catggtgatt gtggaattct gcaaatttgg aaacctgtcc 540

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